	IN 28 TOTAL TRADERS		
	475.18 P>3.00	WI-22211	Homo sapiens full length insert cDNA clone
	475.18 P2.08	stSG48093	ESTs
	475.18 *	A004P27	ESTs, Weakly similar to MULTIDRUG RESI
	475.35 P2.10	stSG9904	ESTs
	475.40 P0.45	sts-AA024696	ESTs _
	475.51 P>3.00	stSG53793	ESTs
,	476.10 P>3.00	Bda98d05	Homo sapiens full length insert cDNA clone
	476.21 P>3.00	sts-H24468	ESTs
	476.21 P>3.00	sts-N94741	ESTs
	476.64 P0.28	stSG22488	ESTs
	476.85 P0.36	stSG44909	ESTs
	477.06 P0.10	stSG54797	ESTs
	477.27 P1.33	stSG48099	ESTs
	477.37 P0.09 *	sts-AA028894	Homo sapiens silencing mediator of retinoic
	477.80 P1.44	stSG52727	EST
	477.80 "	U44799	Human U1-snRNP binding protein homolog mR
	477.80 *	WI-15963	ESTs
	477.80 "	stSG53886	ESTs, Weakly similar to neuronal thread pr
	478.74 P0,01	WIAF-364	ESTs
	479.01 P0.21	WI-21080	ESTs
	479.13 P0.19	A009B29	ESTs
	479.33 P0.22	A006F32 EIF	eukaryotic translation initiation factor 2B
	479.33 P0.19	WIAF-449 EIF	•
	479.33 P0.19 *	WI-15890	H.sapiens mRNA for transmembrane protein r
	479.55 P0.20 *	stSG349	H.sapiens mRNA for transmembrane protein r
;	479.55 " *		IRB2 bradykinin receptor B2
	479.55 "	stSG42540	ESTs
	479.55 "	sts-N26791	ESTs
	479.55 "	stSG53943	ESTs
145.7	479.55 "	stSG49468	EST
A7J.1	479.74 P0.16 481.46 P0.00		Microsatellite marker AFM294ze9
147.5 =		sts-AA007694	EST
	Next interval down	AFM294xg1 D12	S340 Microsatellite anchor marker AFM294xg1
`			

FIG. 2I

	483.87 P0.83		stSG47707		ESTs
l	484.70 P0.93		stSG4060		ESTs
I	484.70 "		stSG62390	GTF2H3	general transcription factor IIH, polypepti
١	484.70 "		stSG42994		ESTs
I	484.73 P0.74		stSG46906		ESTs
	484.80 P0.91		A004X33		ESTs
l	484.91 P1.11		stSG3211		ESTs, Weakly similar to B-cell growth fact
	484.91 "	*	sts-Z41302	BDKRB2	bradykinin receptor B2
١	484.91 "	*	sts-Z41302	BDKRB2	bradykinin receptor B2
I	484.91 "		sts-T58259		ESTs, Weakly similar to B-cell growth fact
	484.91 "		stSG52737		ESTs
I	484.91 "		Bda03b10	UBC	ubiquitin C
l	484.91 "		stSG1936	CD36L1	CD36 antigen (collagen type I receptor, thr
I	484.91 "		sts-AA017225		ESTs
	484.91 P1.15		WI-12212		ESTs
	485.12 P1.18		A004F14		ESTs
l	485.12 P1.18		SGC31333		ESTs
١	485.23 P1.21	*	WI-12482	BDKRB2	bradykinin receptor B2
١	485.23 P1.07		sts-AA017698		ESTs
	485.33 P1.22		WI-12422		ESTs
١	485.51 P1.18		stSG42398		EST
l	485.64 P1.04		sts-AA009669		ESTs
١	486.07 P2.50		stSG21539		EST
	486.13 P1.44		WI-12439		EST
1	486.34 P1.26		sts-W31616	UBC	ubiquitin C
	486.38 P>3.00		stSG54715		ESTs
	486.76 P1.64	×	WI-6921		H.sapiens mRNA for transmembrane protein $r.$.
-	487.08 P>3.00		WI-13120		Human mRNA for KIAA0318 gene, partial cds
	487.23 P>3.00		stSG54353		ESTs
Ì	487.23 P>3.00		stSG22703		EST
	487.28 P>3.00		stSG62698		ESTs
	487.28 P>3.00	*	sts-D60472		Homo sapiens silencing mediator of retinoic
	487.28 P>3.00		stSG36097		ESTs
	487.33 P1.36		sts-U37146		Homo sapiens silencing mediator of retinoic

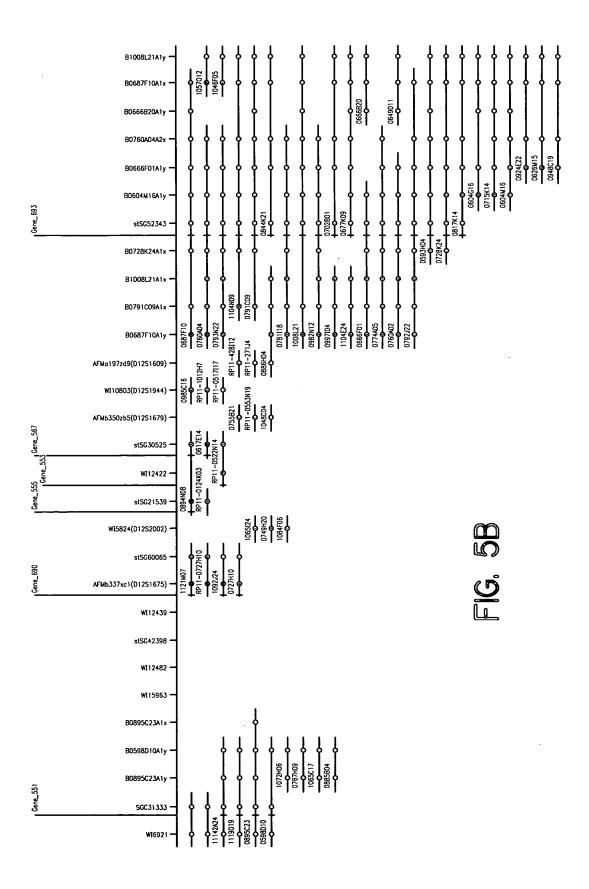
FIG. 2K

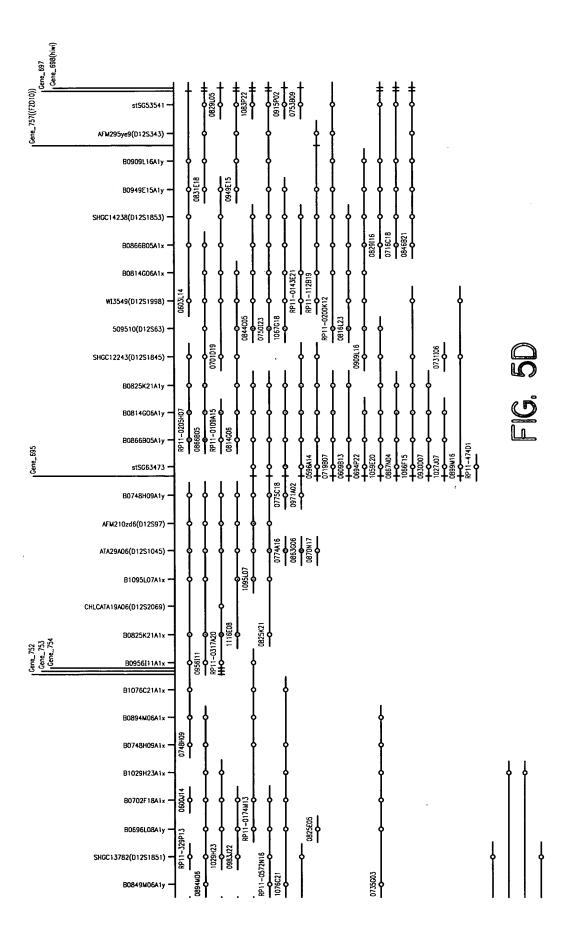
	497.60 P. 2.00				
	487.50 P>3.00		stSG9807		ESTs
	487.50 P>3.00		stSG15434		ESTs
	487.60 P>3.00		stSG53251		ESTs
	487.60 P>3.00		stSG30525	SRRP129	SC35-interacting protein 1
	487.60 P>3.00		stSG46424		ESTs
	487.70 P>3.00		A007A34		ESTs
154.4	487.75 P2.00		AFMa197zd9	D12S1609	Microsatellite marker AFMa197zd9
ı	487.75 P2.02		A006D44		ESTs
	487.80 P>3.00		SGC30248		ESTs, Weakly similar to peptide/histidine
	488.07 P1.68		stSG6320		Homo sapiens clone 24617 mRNA sequence
	488.07 P1.66		stSG6305		Homo sapiens clone 24790 mRNA sequence
	488.07 P0.02		sts-N20163		Homo sapiens full length insert cDNA clone
	488.12 P>3.00		stSG60065		ESTs
	488.12 P>3.00		stSG47723		ESTs
	488.44 P1.59		stSG3292	•	Homo sapiens clone 24790 mRNA sequence
	488.44 P0.03		WIAF-856		EST, Weakly similar to reverse transcripta
	488.65 P1.54		WI-12272		Homo sapiens clone 24790 mRNA sequence
	488.65 P1.82		stSG52343		ESTs
	488.82 P1.80		stSG16387	CPN2	carboxypeptidase N, polypeptide 2, 83kD
	488.97 P1.80		SGC31722		ESTs
	489.07 P0.06		stSG54325		ESTs
	489.07 P>3.00		stSG63473		ESTs
160.9	♦ 489.07 P>3.00		AFMa123xe1	D12S367	Microsatellite marker AFMa123xe1
	489.14 PO.17		sts-T81113		ESTs
	489.29 P0.05		sts-AA025438		EST
	489.50 P1.37	***	Cdalad08		ESTs
	489.50 P0.05		WI-15018		ESTs
	489.50 P1.50		WI-18492		ESTs
	489.57 P1.48		WI-16177		Homo sapiens androgen receptor associated p
,	489.67 P1.44		stSG53307		ESTs
1	489.71 P1.43		stSG53541		Homo sapiens hiwi mRNA, partial cds
	489.71 P1.43		stSG9546		Homo sapiens clone 24617 mRNA sequence
	489.89 P1.56		A006O16		ESTs
	490.10 P1.42		H64839		EST

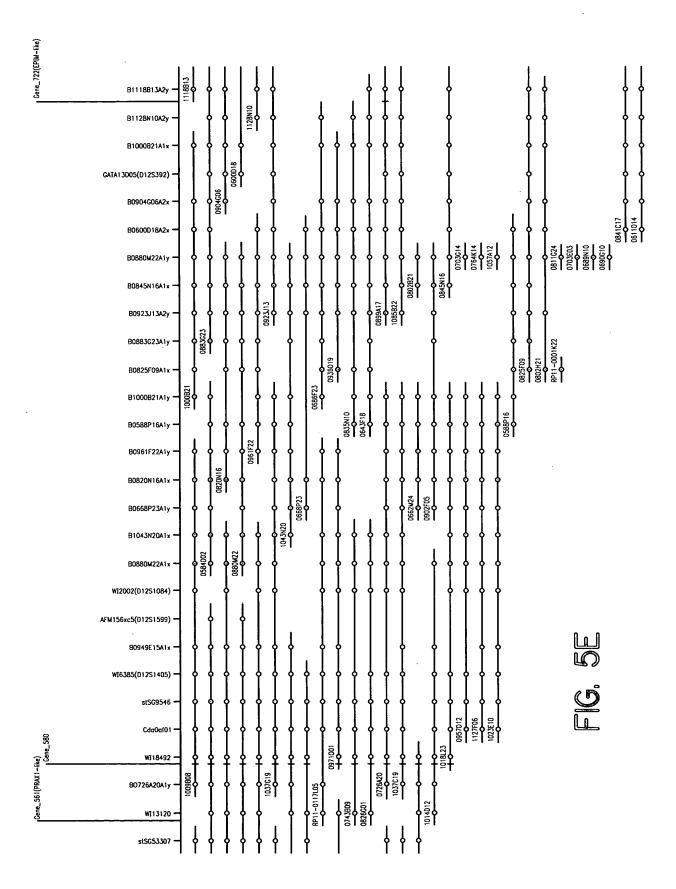
FIG. 2L

499.71	P1.72	stSG30906		ESTs
499.71		stSG43796	MMP17	matrix metalloproteiriase 17 (membrane-insert
499.71	P1.71	sts-X89576	MMP17	matrix metalloproteinase 17 (membrane-insert
499.92	P>3.00	stSG43769		ESTs
500.50	P1.88	stSG26056		EST ₅
500.50	P2.33	SGC30786	KIAA0331	KIAA0331 gene product
500.61	P>3.00	stSG1702		Homo sapiens CAGH32 mRNA, partial cds
500.61	•	sts-N59820		EST _{\$}
500.61	•	stSG42115	KIAA0331	KIAA0331 gene product
500.61	•	IB2452	ULK1	unc-51 (C. elegans)-like kinase 1
500.61	¢1	stSG52521		EST _s
500.61	н	FB9F8		ESTs, Weakly similar to PUTATIVE ATP-D
500.61	11	AA252357		ESTs
500.61	11	stSG4720		Homo sapiens pseudouridine synthase 1 (PUS
500.61	**	sts-AA001424	KIAA0331	KIAA0331 gene product
500.61	P>3.00	stSG31443		ESTs
500.61	P>3.00	stSG49622	ULK1	unc-51 (C. elegans)-like kinase 1
500.61	P2.49	stSG50559		ESTs
501.04	P1.10	stSG54842		ESTs
501.04	P2.03	A008Y05		Unknown
501.89	P2.18	stSG39493		Homo sapiens CAGH32 mRNA, partial cds
501.99	P>3.00	A002A44		Homo sapiens CAGH32 mRNA, partial cds
501.99	P>3.00	sts-H94865	•	EST
501.99	P>3.00	R50113		ESTs
502.10	P1.75	stSG48386		ESTs
502.10	**	stSG50504		ESTs
502.63	P0.06	A006R19		ESTs
502.63	P1.06	WIAF-864		ESTs
502.94	P1.51	stSG54813		ESTs, Weakly similar to peroxisome membran
503.04	P1.42	A004B47		ESTs, Highly similar to DNA polymerase ep
503.25	P0.28	stSG27206		ESTs
503.25	n	stSG40199		Homo sapiens mRNA for KIAA0692 protein, p
503.46	P0.23	stSG8935		ESTs
504.68	P0.69	stSG4731		Homo sapiens mRNA for KIAA0692 protein, p

FIG. 20







0794A21 076B112 0723P10 B0723P10A1y B1056C02A2x Gene_694 stSG54325 stSG42115 B0839D11A1y 987 B1128L12A1y Gene_751 W16077(D12S1322) B0716I10A2x SGC34088 Gene_575 WIAF856 RP11-0579G13 B0610N03A1x 11281.12 B0700A09A2y RP11-0243M05 B1118B13A2x 1083H10 1093F08 0839011 0909E24 B0909E24A1y B0600D18A2y B1128N10A2x B0841C17A2x B0611014A2x Gene_581(TC4, RAN) stSG3292 B0923J13A2x stSG3357 Gene_585 WI16177 1 Gene_584 H64839 B0700A09A2x W112272 Cda0ca07

